
Exemption #5

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National Center for Environmental Assessment
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Washington, DC

From: DeSantis, Joe

Sent: Monday, July 20, 2015 10:29 AM

To: White, Paul; Jones, Samantha; Flowers, Lynn; Walker, Teneille; Callan, Richard; Ilacqua, Vito

Cc: Perovich, Gina; Cogliano, Vincent; Blaine, Susan; bryan.luukinen@icfi.com

Subject: Comments on Nominees? - EPI

Good morning –



Exemption #5

D.

Person you are Nominating as an Expert	Nominee's Affiliation	Nominee's Email Address	Summary of Nominee's
Holly Porter Morgan Nominated by: Laura Hoffman, Barge Park Pals, Newton Creek Alliance	LaGuardia Community College	LaGuardia Community College	<p>As a life long resident of Greenpoint, who entire family has been impacted by sew more, I've been very active in the community. Holly Porter-Morgan is very much av understanding of the cumulative impacts here. And she has been an asset to the N Assistant Professor of Biology with a PhD in Biology. She specializes in Ecology and the New York University for a Masters Degree in Interdisciplinary Studies. She then geospatial technology for conservation as a research scientist at the New York Bot</p> <p>Dr.Porter-Morgan directs the Environmental Science Program at LaGuardia and tea Science. She is currently serving as a research mentor for the NSF-funded Louis Sto</p> <p>Dr. Porter-Morgan's research interests lie at the confluence of Environmental Scie she is studying the potential for bioremediation to supplement proposals for wate Queens. This research examines measures of organismal diversity and water qualiti including the Newtown Creek Alliance.</p> <p>The second portion of her research is a geospatial (GISc) investigation of the enviro surrounding Newtown Creek. This work incorporates field data collected by studen organizations including the EPA, DOH, and DEC.</p>
Dr. Reza Rasoulpour Nominated by: Kamin Johnson, DOW	The Dow Chemical Company	rrasoulpour@dow.com	<p>Dr. Rasoulpour has expertise in performing and interpreting experiments involving published recent peer-reviewed articles on the use of epigenetic endpoints in toxic epigenetic and apical endpoints. He brings an important perspective on the role of</p>
Self	The Dow Chemical Company	kjohnson2@dow.com	<p>I have extensive experience in developmental and reproductive toxicity in both an the mechanisms and outcomes of fetal phthalate exposure and employed a variety have been a lead scientist in an industrial setting interpreting a variety of DART gu mode of action perspective and a potential use standpoint in guideline-based toxic</p>

Janine LaSalle
Nominated by:
 Melissa Rose, UC
 Davis Center for
 Children's
 Environmental
 Health

University of
 Californi, Davis

jmlasalle@ucdavis.edu

Dr. LaSalle's research contributes to understanding the interplay of genetics, environment, and human health. Her lab takes a "Rosetta's stone" approach to decoding the elusive pathways disrupted in rare genetic disorders on the autism spectrum. The laboratory is studying a protein that binds to methylated DNA, methyl CpG binding protein 2 (MeCP2). The syndrome and other neurodevelopmental disorders. In addition, Dr. LaSalle is interested in the Prader-Willi locus that are expressed in postnatal neurons. She is also investigating DNA methylation and chromatin organization in 15q11-13 duplication syndrome. Relevant publications include:

LaSalle JM. 2011. A genomic point-of-view on environmental factors influencing the autism spectrum. *Hum. Mol. Genet.* 20:115-125.

Woods R, Vallero RO, Golub M, Suarez JK, Ta TA, Yasui DH, Chi L-H, Kostyniak PJ, Pessah IN. 2010. Interactions between perinatal PBDE exposure and Mecp2308 mutation. *Hum. Mol. Genet.* 19:115-125.

Mitchell MM, Woods R, Chi L-H, Schmidt RJ, Pessah IN, Kostyniak PJ, LaSalle JM. 2010. Postmortem brain reveal possible environmental epigenetic involvement in 15q11-q13 duplication syndrome. *Mutagenesis*, 33:589-598.

LaSalle JM. 2013. Epigenomic strategies at the interface of genetic and environmental factors. *Hum. Mol. Genet.* 22:115-125.

Self

US Army
 Engineer
 Research and
 Development
 Center

Lyle.D.Burgoon@usace.army.mil

Dr. Burgoon is an expert on the use of epigenetics information in risk screening and assessment (EPA/NCEA), created the first, novel approach to utilizing a data science approach to integrate GWAS information to perform geographic-based risk screening for environmental health. (<http://journals.plos.org/plosone/article?id=10.1371/journal.pone.0121855>).

Dr. Burgoon is a data scientist, risk assessor, and bioinformatician that specializes in gene expression modeling. Dr. Burgoon has published numerous studies on gene expression regulation of the aryl hydrocarbon receptor. In addition, Dr. Burgoon has published on the regulation and placement and how that impacts gene expression (with collaborator Dr. David Schuster) following chemical exposure with Dr. Jay Goodman.

In addition to a strong biological understanding, Dr. Burgoon also provides a statistical analysis. In addition, Dr. Burgoon designs, develops, releases, and reviews bioinformatics software for gene expression, and metabolomics.

Finally, Dr. Burgoon is a former NCEA risk assessor, and one of the lead editors and reviewers of the next generation risk assessment, and leads military science and risk assessment technologies into military science and risk assessment.

**Dr. Tsu-Fan
 Cheng**

Nominated by:
 Geoffrey Patton,
 US FDA CFSAN

U.S. FDA CFSAN

Tsu-Fan.Cheng@fda.hhs.gov

Dr. Cheng is a toxicology reviewer in the FDA who also researches how epigenetics can be used in risk assessment. His research in epigenetics has resulted in two publications in recent years. The first publication shows that cadmium, arsenic and nickel, etc.) can alter the epigenetic status, and what kind of epigenetics that resulted from the exposure to heavy metals (Cheng et al., 2012). The second publication shows the lymphoma assay (MLA) as a screening tool to identify chemicals that alter epigenetics. Dr. Cheng postulates that with a slight modification to the original protocol of MLA, one can use MLA to identify chemicals that alter epigenetics or both. Dr. Cheng has been an esteemed member of the Division's Genetic Toxicology Committee, and has been leading reviews of testing methods for application to regulatory submissions to the FDA.

**Self –
 Georgio
 Assennato**

Director, Apulia
 Environmental
 Protection
 Agency, Italy

g.assennat@arpa.puglia.it

Health risk and impact assessment of industrial emissions. Author of several publications on the health risk assessment at Brindisi (petrochemical and coal-fired power plant) and Taranto (incinerators).

**Self – John
French**

UNC-Chapel Hill jfren43@email.unc.edu

Dr. French continues to focus on the development of new population-based approaches to study individual variation in toxic response to environmental exposures and to identify genetic and epigenetic variants associated with the development of disease. He has proposed that both genetic and epigenetic variant identification be used to assess the potential for reverse genetics validation of the variants with the goal of understanding the role of the recent Environ Health Perspect. 2015 Mar;123(3):237-45. P. French has the potential to perform GWAS (and EWAS) in a genetically-diverse mouse population and epigenetic diversity observed in the human population to aid in understanding disease toxicity.

Dr. French earned his Ph.D. at North Carolina State University and worked for the US Food and Drug Administration, National Toxicology Program, and the NIEHS. He has organized and participated in a number of national and international meetings and has published more than 120 peer-reviewed papers. During his tenure at NTP/NIEHS, he was involved in leadership and research as Group Leader, Transgenic Carcinogenesis and Susceptibility Initiative from 2007-2013. The NTP Host Susceptibility Initiative is a testing program created to increase understanding of the genetic and environmental factors that make some individuals more susceptible to toxicants and/or disease susceptibility. The initiative has aided understanding of how some environments are hazardous to some individuals and not to others. Dr. French is a Special Volunteer working on data mining from completed projects and is currently working on mouse models in toxicology and environmental health. Currently he is a member of the Pharmacogenetics and Individualized Therapy and the Department of Environmental Health Sciences at UNC-Chapel Hill.